

IN THE CLAIMS

Please cancel claims 1-28 without prejudice or disclaimer.

Please add new claims 29-47.

This listing of the claims replaces all prior versions of the claims in the application.

1.-28. (Canceled)

29. (New) An isolated polypeptide selected from the group consisting of:

- a) a polypeptide comprising an amino acid sequence of SEQ ID NO:5,
- b) a polypeptide comprising a naturally occurring amino acid sequence at least 90% identical to an amino acid sequence of SEQ ID NO:5,
- c) a biologically active fragment of a polypeptide having an amino acid sequence of SEQ ID NO:5, and
- d) an immunogenic fragment of a polypeptide having an amino acid sequence of SEQ ID NO:5.

30. (New) An isolated polypeptide of claim 29 comprising an amino acid sequence of SEQ ID NO:5.

31. (New) An isolated polynucleotide encoding a polypeptide of claim 29.

32. (New) An isolated polynucleotide encoding a polypeptide of claim 30.

33. (New) An isolated polynucleotide of claim 32 comprising a polynucleotide sequence of SEQ ID NO:13.

34. (New) A recombinant polynucleotide comprising a promoter sequence operably linked to a polynucleotide of claim 31.

35. (New) A cell transformed with a recombinant polynucleotide of claim 34.
36. (New) A method of producing a polypeptide of claim 29, the method comprising:
- a) culturing a cell under conditions suitable for expression of the polypeptide, wherein said cell is transformed with a recombinant polynucleotide, and said recombinant polynucleotide comprises a promoter sequence operably linked to a polynucleotide encoding the polypeptide of claim 29, and
 - b) recovering the polypeptide so expressed.
37. (New) A method of claim 36, wherein the polypeptide comprises an amino acid sequence of SEQ ID NO:5.
38. (New) An isolated antibody which specifically binds to a polypeptide of claim 29.
39. (New) An isolated polynucleotide selected from the group consisting of:
- a) a polynucleotide comprising a polynucleotide sequence of SEQ ID NO:13,
 - b) a polynucleotide comprising a naturally occurring polynucleotide sequence at least 90% identical to a polynucleotide sequence of SEQ ID NO:13,
 - c) a polynucleotide complementary to a polynucleotide of a),
 - d) a polynucleotide complementary to a polynucleotide of b), and
 - e) an RNA equivalent of a)-d).
40. (New) A method of detecting a target polynucleotide in a sample, said target polynucleotide having a sequence of a polynucleotide of claim 39, the method comprising:
- a) hybridizing the sample with a probe comprising at least 20 contiguous nucleotides comprising a sequence complementary to said target polynucleotide in the sample, and which probe specifically hybridizes to said target polynucleotide, under conditions

whereby a hybridization complex is formed between said probe and said target polynucleotide or fragments thereof, and

- b) detecting the presence or absence of said hybridization complex, and, optionally, if present, the amount thereof.

41. (New) A method of detecting a target polynucleotide in a sample, said target polynucleotide having a sequence of a polynucleotide of claim 39, the method comprising:

- a) amplifying said target polynucleotide or fragment thereof using polymerase chain reaction amplification, and
- b) detecting the presence or absence of said amplified target polynucleotide or fragment thereof, and, optionally, if present, the amount thereof.

42. (New) A composition comprising a polypeptide of claim 29 and a pharmaceutically acceptable excipient.

43. (New) A composition of claim 42, wherein the polypeptide comprises an amino acid sequence of SEQ ID NO:5.

44. (New) A method of screening for a compound that specifically binds to the polypeptide of claim 29, the method comprising:

- a) combining the polypeptide of claim 29 with at least one test compound under suitable conditions, and
- b) detecting binding of the polypeptide of claim 29 to the test compound, thereby identifying a compound that specifically binds to the polypeptide of claim 29.

45. (New) A method of screening for a compound that modulates the activity of the polypeptide of claim 29, the method comprising:

- a) combining the polypeptide of claim 29 with at least one test compound under conditions permissive for the activity of the polypeptide of claim 29,
- b) assessing the activity of the polypeptide of claim 29 in the presence of the test compound, and
- c) comparing the activity of the polypeptide of claim 29 in the presence of the test compound with the activity of the polypeptide of claim 29 in the absence of the test compound, wherein a change in the activity of the polypeptide of claim 29 in the presence of the test compound is indicative of a compound that modulates the activity of the polypeptide of claim 29.

46. (New) A method of screening a compound for effectiveness in altering expression of a target polynucleotide, wherein said target polynucleotide comprises a sequence of claim 33, the method comprising:

- a) contacting a sample comprising the target polynucleotide with a compound, under conditions suitable for the expression of the target polynucleotide,
- b) detecting altered expression of the target polynucleotide, and
- c) comparing the expression of the target polynucleotide in the presence of varying amounts of the compound and in the absence of the compound.

47. (New) A method of screening for potential toxicity of a test compound, the method comprising:

- a) treating a biological sample containing nucleic acids with the test compound,
- b) hybridizing the nucleic acids of the treated biological sample with a probe comprising at least 20 contiguous nucleotides of a polynucleotide of claim 39 under conditions whereby a specific hybridization complex is formed between said probe and a target polynucleotide in the biological sample, said target polynucleotide comprising a polynucleotide sequence of a polynucleotide of claim 39 or fragment thereof,
- c) quantifying the amount of hybridization complex, and
- d) comparing the amount of hybridization complex in the treated biological sample with the amount of hybridization complex in an untreated biological sample, wherein a difference in the amount of hybridization complex in the treated biological sample indicates potential toxicity of the test compound.